

## SEQUENCE LISTING

<110> FAGAN, Richard Joseph  
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 BOSCHERT, Ursula  
 CHVATCHKO, Yolande

<120> CYTOKINE ANTAGONIST MOLECULES

<130> C.R.116

<140> US 10/579,113  
 <141> 2006-05-11

<150> PCT/GB2004/004772  
 <151> 2004-11-12

<150> GB0326393.6  
 <151> 2003-11-12

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 <212> PRT  
 <213> Homo sapiens

<400> 2  
 Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg  
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 Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp  
 20 25

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ggcagctgaa gggggacaag ccagtgaccg tgggtgcagtc cattggcaca gaggtcatcg 180
gcaccctgcg gcttgactat cgagaccgta tccgactctt tgaaaatggc tccctgcttc 240
tcagcgacct gcagctggcc gatgagggca cctatgaggt cgagatctcc atcaccgacg 300
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<210> 4
<211> 114
<212> PRT
<213> Homo sapiens

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20 25 30
Ser Asp Arg Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val
35 40 45
Thr Val Val Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro
50 55 60
Asp Tyr Arg Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu
65 70 75 80
Ser Asp Leu Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser
85 90 95
Ile Thr Asp Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val
100 105 110
Asp Val

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<210> 5
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<212> DNA
<213> Homo sapiens

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tgcccatttc gagggcacag gtgttggtgg cttcaaccac tgtgctggag ctcagcgagg 60
ccttcacctt gaactgctca catgagaatg gcaccaagcc cagctacacc tggctgaagg 120
atggcaagcc cctcctcaat gactcgagaa tgctcctgtc ccccgaccaa aagggtgctca 180
ccatcaccgg cgtgctcatg gaggatgacg acctgtacag ctgcatggtg gagaacccca 240
tcagccaggg ccgcagcctg cctgtcaaga tcaccgtata ca 282

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<210> 6
<211> 94
<212> PRT
<213> Homo sapiens

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Pro Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu
1 5 10 15

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Leu Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys  
                   20                  25                  30

Pro Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser  
           35                  40                  45

Arg Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val  
       50                  55                  60

Leu Met Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile  
   65                  70                  75                  80

Ser Gln Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg  
                   85                  90

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 <212> DNA  
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 tgacagtctg tgccctgctgg aaaccctcca aaag 94

<210> 8  
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 <213> Homo sapiens

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Val Thr Leu Val Thr Val Cys Ala Cys Trp Lys Pro Ser Lys Arg  
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<210> 9  
 <211> 74  
 <212> DNA  
 <213> Homo sapiens

<400> 9  
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 cctgaaacca gaag 74

<210> 10  
 <211> 25  
 <212> PRT  
 <213> Homo sapiens

<400> 10  
 Lys Gln Lys Lys Leu Glu Lys Gln Asn Ser Leu Glu Tyr Met Asp Gln  
 1                  5                  10                  15

Asn Asp Asp Arg Leu Lys Pro Glu Ala  
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 <211> 71  
 <212> DNA  
 <213> Homo sapiens

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 <212> PRT  
 <213> Homo sapiens

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 Asp Thr Leu Pro Arg Ser Gly Glu Gln Glu Arg Lys Asn Pro Met Ala  
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Leu Tyr Ile Leu Lys Asp Lys  
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<210> 13  
 <211> 303  
 <212> DNA  
 <213> Homo sapiens

<400> 13  
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 tctgccccgc gctacccgcg ctccccagcg cgctccccag ccaccggccg gacacactcg 180  
 tcgcccgcga gggccccgag ctgcgccggc cgctcgcgca gcgcctcgcg cacactgcgg 240  
 actgcggggc tgcacataat ccgcgagcaa gacgaggccg gcccgggtgga gatcagcgcc 300  
 tga 303

<210> 14  
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 <212> PRT  
 <213> Homo sapiens

<400> 14  
 Asp Ser Pro Glu Thr Glu Glu Asn Pro Ala Pro Glu Pro Arg Ser Ala  
 1                  5                  10                  15

Thr Glu Pro Gly Pro Pro Gly Tyr Ser Val Ser Pro Ala Val Pro Gly  
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Arg Ser Pro Gly Leu Pro Ile Arg Ser Ala Arg Arg Tyr Pro Arg Ser  
                   35                  40                  45

Pro Ala Arg Ser Pro Ala Thr Gly Arg Thr His Ser Ser Pro Pro Arg  
                   50                  55                  60

Ala Pro Ser Ser Pro Gly Arg Ser Arg Ser Ala Ser Arg Thr Leu Arg  
65 70 75 80

Thr Ala Gly Val His Ile Ile Arg Glu Gln Asp Glu Ala Gly Pro Val  
85 90 95

Glu Ile Ser Ala  
100

<210> 15  
<211> 1251  
<212> DNA  
<213> Homo sapiens

<400> 15  
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cgctgatcc atggcacctg ggggaagtgc gctctgcttt ctgtgcagta cagcagtacc 180  
agcagcgaca ggctgtagt gaagtggcag ctgaagcggg acaagccagt gaccgtggtg 240  
cagtccattg gcacagaggt catcggcacc ctgcggcctg actatcgaga ccgtatccga 300  
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gaggtcgaga tctccatcac cgacgacacc ttcaactggg agaagacat caaccttact 420  
gtagatgtgc ccatttcgag gccacaggtg ttggtggctt caaccactgt gctggagctc 480  
agcgaggcct tcaccttgaa ctgctcacat gagaatggca ccaagcccag ctacacctgg 540  
ctgaaggatg gcaagcccct cctcaatgac tcgagaatgc tcctgtcccc cgaccaaag 600  
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aaccatca gccagggcgc cagcctgcct gtcaagatca ccgtatacag aagaagctcc 720  
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gcctgctgga aacctccaa aaggaaacag aagaagctag aaaagcaaaa ctccctggaa 840  
tacatggatc agaattgatga ccgcctgaaa ccagaagcag acacctccc tcgaagtggg 900  
gagcaggaac ggaagaacct catggcactc tatatcctga aggacaagga ctccccggag 960  
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<210> 16  
<211> 416  
<212> PRT  
<213> Homo sapiens

<400> 16  
Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg  
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Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu  
20 25 30

Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly  
35 40 45

Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg  
50 55 60

Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val  
 65 70 75 80  
 Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg  
 85 90 95  
 Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu  
 100 105 110  
 Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp  
 115 120 125  
 Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro  
 130 135 140  
 Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu  
 145 150 155 160  
 Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro  
 165 170 175  
 Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg  
 180 185 190  
 Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu  
 195 200 205  
 Met Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser  
 210 215 220  
 Gln Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser  
 225 230 235 240  
 Leu Tyr Ile Ile Leu Ser Thr Gly Gly Ile Phe Leu Leu Val Thr Leu  
 245 250 255  
 Val Thr Val Cys Ala Cys Trp Lys Pro Ser Lys Arg Lys Gln Lys Lys  
 260 265 270  
 Leu Glu Lys Gln Asn Ser Leu Glu Tyr Met Asp Gln Asn Asp Asp Arg  
 275 280 285  
 Leu Lys Pro Glu Ala Asp Thr Leu Pro Arg Ser Gly Glu Gln Glu Arg  
 290 295 300  
 Lys Asn Pro Met Ala Leu Tyr Ile Leu Lys Asp Lys Asp Ser Pro Glu  
 305 310 315 320  
 Thr Glu Glu Asn Pro Ala Pro Glu Pro Arg Ser Ala Thr Glu Pro Gly  
 325 330 335  
 Pro Pro Gly Tyr Ser Val Ser Pro Ala Val Pro Gly Arg Ser Pro Gly  
 340 345 350  
 Leu Pro Ile Arg Ser Ala Arg Arg Tyr Pro Arg Ser Pro Ala Arg Ser

355                                      360                                      365  
 Pro Ala Thr Gly Arg Thr His Ser Ser Pro Pro Arg Ala Pro Ser Ser  
       370                                      375                                      380  
 Pro Gly Arg Ser Arg Ser Ala Ser Arg Thr Leu Arg Thr Ala Gly Val  
 385                                      390                                      395                                      400  
 His Ile Ile Arg Glu Gln Asp Glu Ala Gly Pro Val Glu Ile Ser Ala  
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<210> 17  
 <211> 1257  
 <212> DNA  
 <213> Mus musculus

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 cgtctgatcc acggcacagt ggggaagtgc gccctgcttt ccgtgcagta cagtagcacc 180  
 agcagcgaca agcccgtggt gaagtggcag ctgaagcgtg acaagccagt gaccgtggtg 240  
 cagtctatag gcacagaggt cattggcact ctgcggcctg actatcgaga ccgtatccgg 300  
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 gtggatgtgc ccatttcaag gccgcaggta ttagtggctt caacctactgt gctggagctc 480  
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 gtgctcacca tcaccgagt actcatggaa gatgacgacc tgtacagctg tgtgggtggag 660  
 aaccccatca gccaggtcct cagcctgcct gtcaagatca ctgtgtatag aagaagctcc 720  
 ctctatatca tcttgtctac aggaggcact ttctccttgg tgacctggt gacagtttgt 780  
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 agtggagaac aggagcggaa gaacccaatg gcactctata tcctgaagga taaggattcc 960  
 tcagagccag atgaaaaccc tgctacagag ccacggagca ccacagaacc cggccccct 1020  
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 cgcgctacc cgcgctcccc agcacgttcc cctgccactg gccggacgca cacgtcgcca 1140  
 ccgcggggcc cgagctcgcc aggcgctcg cgcagctctt cgcgctcact gcggactgca 1200  
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<210> 18  
 <211> 418  
 <212> PRT  
 <213> Mus musculus

<400> 18  
 Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg  
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                                     20                                      25                                      30  
 Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly  
                                     35                                      40                                      45

Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Lys  
 50 55 60  
 Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val  
 65 70 75 80  
 Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg  
 85 90 95  
 Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu  
 100 105 110  
 Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp  
 115 120 125  
 Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro  
 130 135 140  
 Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu  
 145 150 155 160  
 Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro  
 165 170 175  
 Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg  
 180 185 190  
 Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu  
 195 200 205  
 Met Glu Asp Asp Asp Leu Tyr Ser Cys Val Val Glu Asn Pro Ile Ser  
 210 215 220  
 Gln Val Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser  
 225 230 235 240  
 Leu Tyr Ile Ile Leu Ser Thr Gly Gly Ile Phe Leu Leu Val Thr Leu  
 245 250 255  
 Val Thr Val Cys Ala Cys Trp Lys Pro Ser Lys Lys Ser Arg Lys Lys  
 260 265 270  
 Arg Lys Leu Glu Lys Gln Asn Ser Leu Glu Tyr Met Asp Gln Asn Asp  
 275 280 285  
 Asp Arg Leu Lys Ser Glu Ala Asp Thr Leu Pro Arg Ser Gly Glu Gln  
 290 295 300  
 Glu Arg Lys Asn Pro Met Ala Leu Tyr Ile Leu Lys Asp Lys Asp Ser  
 305 310 315 320  
 Ser Glu Pro Asp Glu Asn Pro Ala Thr Glu Pro Arg Ser Thr Thr Glu  
 325 330 335  
 Pro Gly Pro Pro Gly Tyr Ser Val Ser Pro Pro Val Pro Gly Arg Ser



Ser Ala

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cgctgatcc	atggcaccgt	ggggaagtgc	gctctgcttt	ctgtgcagta	cagcagtacc		180
agcagcgaca	ggcctgtagt	gaagtggcag	ctgaagcggg	acaagccagt	gaccgtggtg		240
cagtccattg	gcacagagggt	catcggcacc	ctgcggcctg	actatcgaga	ccgtatccga		300
ctctttgaaa	atggctccct	gctttctcagc	gacctgcagc	tggccgatga	gggcacctat		360
gaggctcgaa	tctccctcac	cgacgcagacc	ttcactgggg	agaagaccat	caaccttact		420
gtagatgtgc	ccatttcag	gccacaggtg	ttggtggctt	caaccactgt	gctggagctc		480
agcagggcct	tcaccttgaa	ctgtccagat	gagaatggca	ccaagcccag	ctacaccttg		540
ctgaaggatg	gcaagccctt	cctcaatgac	tcgagaatgc	tcttgtcccc	cgacaaaaag		600
gtgtcacca	tcaccgcgt	gctcatggag	gatgacgacc	tgtacagctg	catggtggag		660
aaccccatca	gccagggccg	cagcctgcct	gtcaagatca	ccgtatacag	aagaagctcc		720

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<400>      20
Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg
1              5              10              15

Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu
              20              25              30

Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly
              35              40              45

Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg
              50              55              60

Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val

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65		70		75		80
Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg						
	85			90		95
Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu						
	100			105		110
Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp						
	115			120		125
Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro						
	130			135		140
Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu						
	145			150		155
Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro						
	165			170		175
Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg						
	180			185		190
Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu						
	195			200		205
Met Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser						
	210			215		220
Gln Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser						
	225			230		235
						240

<210> 21  
 <211> 621  
 <212> DNA  
 <213> Homo sapiens

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 gacaagccag tgaccgtggg gcagtcatt ggcacagagg tcatcggcac cctgcggcct 180  
 gactatcgag accgtatccg actctttgaa aatggctccc tgcttctcag cgacctgcag 240  
 ctggccgatg agggcaccta tgaggtcgag atctccatca ccgacgacac cttoactggg 300  
 gagaagacca tcaaccttac tgtagatgtg ccattttcga ggccacaggt gttgggtggc 360  
 tcaaccactg tgctggagct cagcgaggcc ttacacctga actgttcaca tgagaatggc 420  
 accaagccca gctacacctg gctgaaggat ggcaagcccc tcttcaatga ctcgagaatg 480  
 ctctgtctcc ccgaccacaaa ggtgctcacc atcaccgcgc tgctcatgga ggatgacgac 540  
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<210> 22  
 <211> 207  
 <212> PRT  
 <213> Homo sapiens

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<400>    22
Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly Lys
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Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg Pro
          20          25          30

Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val Gln
          35          40          45

Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg Asp
50          55          60

Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu Gln
65          70          75          80

Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp Asp
          85          90          95

Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro Ile
          100          105          110

Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu Ser
          115          120          125

Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro Ser
          130          135          140

Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg Met
145          150          155          160

Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu Met
          165          170          175

Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser Gln
          180          185          190

Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser
          195          200          205

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<210>    23
<211>    328
<212>    DNA
<213>    Homo sapiens

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gacaagccag tgaccgtggt gcagtcattt ggcacagagg tcatcggcac cctgcggcct    180
gactatcgag accgtatccg actctttgaa aatggctccc tgcttctcag cgacctgcag    240
ctggccgatg agggcaccta tgaggtcgag atctccatca ccgacgacac cttcactggg    300
gagaagacca tcaaccttac tgtagatg                                     328

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<210>    24

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<211> 110  
 <212> PRT  
 <213> Homo sapiens

<400> 24  
 Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly Lys  
 1 5 10 15  
 Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg Pro  
 20 25 30  
 Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val Gln  
 35 40 45  
 Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg Asp  
 50 55 60  
 Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu Gln  
 65 70 75 80  
 Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp Asp  
 85 90 95  
 Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val  
 100 105 110

<210> 25  
 <211> 1152  
 <212> DNA  
 <213> Homo sapiens

<400> 25  
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 tctgtgcagt acagcagtag cagcagcgac aggcctgtag tgaagtggca gctgaagcgg 120  
 gacaagccag tgaccgtggt gcagtcattt ggacagagg tcatcggcac cctgcggcct 180  
 gactatcgag accgtatccg actctttgaa aatggctccc tgcttctcag cgacctgcag 240  
 ctggccgatg agggcaccta tgaggtcgag atctccatca ccgacgacac cttcactggg 300  
 gagaagacca tcaaccttac tgtagatgtg cccatttcga ggccacaggt gttgggtggc 360  
 tcaaccactg tgctggagct cagcgaggcc ttacacctga actgctcaca tgagaatggc 420  
 accaagccca gctacacctg gctgaaggat ggcaagcccc tcctcaatga ctcgagaatg 480  
 ctccgtgtcc ccgacccaaa ggtgctcacc atcaccgcgg tgctcatgga ggatgacgac 540  
 ctgtacagct gcatgggtgga gaaccccatc agccagggcc gcagcctgcc tgtcaagatc 600  
 accgtataca gaagaagctc cctttacatc atcttgtcta caggaggcat cttcctcctt 660  
 gtgaccttgg tgacagtctg tgccctgctg aaacctcca aaaggaaaca gaagaagcta 720  
 gaaaagcaaa actccctgga atacatggat cagaatgat accgcctgaa accagaagca 780  
 gacacctcc ctcgaagtgg tgagcaggaa cggaagaacc ccatggcact ctatatcctg 840  
 aaggacaagg actccccgga gaccgaggag aaccgggcc cgagcctcg aagcgcgacg 900  
 gagcccgccc cgcccgcta ctccgtgtct cccgcccgtc ccggccgctc gccggggctg 960  
 cccatccgct ctgcccgccg ctaccgcgcg tccccagcgc gctccccagc caccggccgg 1020  
 acacactcgt cgccgcccag ggccccgagc tcgcccggcc gctcgcgcag cgctcgcgc 1080  
 aactgcgga ctgcgggctg gcacataatc cgcgagcaag acgaggccgg cccgggtggag 1140  
 atcagcgccct ga 1152

<210> 26  
 <211> 383

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<212>   PRT
<213>   Homo sapiens

<400>   26
Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly Lys
1          5          10          15

Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg Pro
          20          25          30

Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val Gln
          35          40          45

Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg Asp
          50          55          60

Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu Gln
          65          70          75          80

Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp Asp
          85          90          95

Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro Ile
          100          105          110

Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu Ser
          115          120          125

Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro Ser
          130          135          140

Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg Met
          145          150          155          160

Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu Met
          165          170          175

Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser Gln
          180          185          190

Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser Leu
          195          200          205

Tyr Ile Ile Leu Ser Thr Gly Gly Ile Phe Leu Leu Val Thr Leu Val
          210          215          220

Thr Val Cys Ala Cys Trp Lys Pro Ser Lys Arg Lys Gln Lys Lys Leu
          225          230          235          240

Glu Lys Gln Asn Ser Leu Glu Tyr Met Asp Gln Asn Asp Asp Arg Leu
          245          250          255

Lys Pro Glu Ala Asp Thr Leu Pro Arg Ser Gly Glu Gln Glu Arg Lys
          260          265          270

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Asn Pro Met Ala Leu Tyr Ile Leu Lys Asp Lys Asp Ser Pro Glu Thr
    275                      280                      285

Glu Glu Asn Pro Ala Pro Glu Pro Arg Ser Ala Thr Glu Pro Gly Pro
    290                      295                      300

Pro Gly Tyr Ser Val Ser Pro Ala Val Pro Gly Arg Ser Pro Gly Leu
305                      310                      315                      320

Pro Ile Arg Ser Ala Arg Arg Tyr Pro Arg Ser Pro Ala Arg Ser Pro
    325                      330                      335

Ala Thr Gly Arg Thr His Ser Ser Pro Pro Arg Ala Pro Ser Ser Pro
    340                      345                      350

Gly Arg Ser Arg Ser Ala Ser Arg Thr Leu Arg Thr Ala Gly Val His
    355                      360                      365

Ile Ile Arg Glu Gln Asp Glu Ala Gly Pro Val Glu Ile Ser Ala
    370                      375                      380

<210> 27
<211> 256
<212> PRT
<213> Homo sapiens

<400> 27
Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg
1      5      10      15

Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu
    20      25      30

Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly
    35      40      45

Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg
    50      55      60

Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val
    65      70      75      80

Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg
    85      90      95

Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu
    100     105     110

Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp
    115     120     125

Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro
    130     135     140

Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu

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```

145              150              155              160
Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro
      165              170              175
Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg
      180              185              190
Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu
      195              200              205
Met Glu Asp Asp Asp Leu Asp Ser Cys Val Val Glu Asn Pro Ile Asn
      210              215              220
Gln Gly Arg Thr Leu Pro Cys Lys Ile Thr Val Tyr Lys Lys Ser Ser
225              230              235              240
Leu Ser Ser Ile Trp Leu Gln Glu Ala Phe Ser Ser Leu Gly Pro Trp
      245              250              255

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<210> 28
<211> 256
<212> PRT
<213> Homo sapiens

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<400> 28
Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg
1      5      10      15
Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu
      20      25      30
Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly
      35      40      45
Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg
      50      55      60
Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val
      65      70      75      80
Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg
      85      90      95
Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu
      100      105      110
Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp
      115      120      125
Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro
      130      135      140
Ile Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu

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```

145                150                155                160
Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro
      165                170                175
Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg
      180                185                190
Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu
      195                200                205
Met Glu Asp Asp Asp Leu Asp Ser Cys Val Val Glu Asn Pro Ile Asn
      210                215                220
Gln Gly Arg Thr Leu Pro Cys Lys Ile Thr Val Tyr Lys Lys Ser Ser
225                230                235                240
Phe Tyr Ile Ile Cys Leu Lys Glu Ala Ser Ser Ser Phe Gly Pro Trp
      245                250                255

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<210> 29
<211> 213
<212> PRT
<213> Homo sapiens

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<400> 29
Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly Lys
1                5                10                15
Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg Pro
      20                25                30
Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val Gln
      35                40                45
Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg Asp
      50                55                60
Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu Gln
      65                70                75                80
Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp Asp
      85                90                95
Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro Ile
      100                105                110
Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu Ser
      115                120                125
Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro Ser
      130                135                140
Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg Met

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145              150              155              160
Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu Met
              165              170              175
Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser Gln
              180              185              190
Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser His
              195              200              205
His His His His His
              210

<210> 30
<211> 439
<212> PRT
<213> Homo sapiens

<400> 30
Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly Lys
1              5              10              15
Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg Pro
              20              25              30
Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val Gln
              35              40              45
Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg Asp
              50              55              60
Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu Gln
65              70              75              80
Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp Asp
              85              90              95
Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro Ile
              100              105              110
Ser Arg Pro Gln Val Leu Val Ala Ser Thr Thr Val Leu Glu Leu Ser
              115              120              125
Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro Ser
              130              135              140
Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg Met
145              150              155              160
Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu Met
              165              170              175
Glu Asp Asp Asp Leu Tyr Ser Cys Met Val Glu Asn Pro Ile Ser Gln
              180              185              190

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Gly Arg Ser Leu Pro Val Lys Ile Thr Val Tyr Arg Arg Ser Ser Glu  
 195 200 205  
 Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro  
 210 215 220  
 Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys  
 225 230 235 240  
 Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val  
 245 250 255  
 Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp  
 260 265 270  
 Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr  
 275 280 285  
 Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp  
 290 295 300  
 Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu  
 305 310 315 320  
 Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg  
 325 330 335  
 Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys  
 340 345 350  
 Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp  
 355 360 365  
 Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys  
 370 375 380  
 Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser  
 385 390 395 400  
 Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser  
 405 410 415  
 Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser  
 420 425 430  
 Leu Ser Leu Ser Pro Gly Lys  
 435  
 <210> 31  
 <211> 186  
 <212> PRT  
 <213> Homo sapiens  
 <400> 31

Val Arg Leu Ile His Gly Thr Val Gly Lys Ser Ala Leu Leu Ser Val  
 1 5 10 15  
 Gln Tyr Ser Ser Thr Ser Ser Asp Arg Pro Val Val Lys Trp Gln Leu  
 20 25 30  
 Lys Arg Asp Lys Pro Val Thr Val Val Gln Ser Ile Gly Thr Glu Val  
 35 40 45  
 Ile Gly Thr Leu Arg Pro Asp Tyr Arg Asp Arg Ile Arg Leu Phe Glu  
 50 55 60  
 Asn Gly Ser Leu Leu Leu Ser Asp Leu Gln Leu Ala Asp Glu Gly Thr  
 65 70 75 80  
 Tyr Glu Val Glu Ile Ser Ile Thr Asp Asp Thr Phe Thr Gly Glu Lys  
 85 90 95  
 Thr Ile Asn Leu Thr Val Asp Val Pro Ile Ser Arg Pro Gln Val Leu  
 100 105 110  
 Val Ala Ser Thr Thr Val Leu Glu Leu Ser Glu Ala Phe Thr Leu Asn  
 115 120 125  
 Cys Ser His Glu Asn Gly Thr Lys Pro Ser Tyr Thr Trp Leu Lys Asp  
 130 135 140  
 Gly Lys Pro Leu Leu Asn Asp Ser Arg Met Leu Leu Ser Pro Asp Gln  
 145 150 155 160  
 Lys Val Leu Thr Ile Thr Arg Val Leu Met Glu Asp Asp Asp Leu Tyr  
 165 170 175  
 Ser Cys Met Val Glu Asn Pro Ile Ser Gln  
 180 185

<210> 32  
 <211> 256  
 <212> PRT  
 <213> Homo sapiens

<400> 32

Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg  
 1 5 10 15  
 Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu  
 20 25 30  
 Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly  
 35 40 45  
 Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg  
 50 55 60

```

Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val
65              70              75              80

Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg
85              90              95

Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu
100            105            110

Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp
115            120            125

Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro
130            135            140

Ile Ser Arg Pro Gln Val Leu Gly Ala Ser Thr Thr Val Leu Glu Leu
145            150            155            160

Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro
165            170            175

Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg
180            185            190

Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu
195            200            205

Met Glu Asp Asp Asp Leu Tyr Ser Cys Val Val Glu Asn Pro Ile Asn
210            215            220

Gln Gly Arg Thr Leu Pro Cys Lys Ile Thr Glu Tyr Arg Lys Ser Ser
225            230            235            240

Leu Ser Ser Ile Trp Leu Gln Glu Ala Phe Ser Ser Leu Gly Pro Trp
245            250            255

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